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GRANT NUMBER DAMD17-96-1-6181

TITLE: Role of Bcl-2 in Breast Cancer Progression

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REPORT DATE: September 1998

TYPE OF REPORT: Annual

PREPARED FOR: Commanding General  
U.S. Army Medical Research and Materiel Command  
Fort Detrick, Maryland 21702-5012

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# REPORT DOCUMENTATION PAGE

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1. AGENCY USE ONLY (Leave blank)		2. REPORT DATE September 1998	3. REPORT TYPE AND DATES COVERED Annual (1 Aug 97 - 31 Jul 98)	
4. TITLE AND SUBTITLE Role of Bcl-2 in Breast Cancer Progression			5. FUNDING NUMBERS DAMD17-96-1-6181	
6. AUTHOR(S) Hyeong-Reh C. Kim, Ph.D.				
7. PERFORMING ORGANIZATION NAME(S) AND ADDRESS(ES) Wayne State University Detroit, Michigan 48202			8. PERFORMING ORGANIZATION REPORT NUMBER	
9. SPONSORING / MONITORING AGENCY NAME(S) AND ADDRESS(ES) U.S. Army Medical Research and Materiel Command Fort Detrick, Maryland 21702-5012			10. SPONSORING / MONITORING AGENCY REPORT NUMBER	
11. SUPPLEMENTARY NOTES				
12a. DISTRIBUTION / AVAILABILITY STATEMENT Approved for Public Release; Distribution Unlimited			12b. DISTRIBUTION CODE	
13. ABSTRACT (Maximum 200 words)  The anti-apoptotic gene bcl-2 is frequently overexpressed in many human tumors including invasive breast cancer. Involvement of bcl-2 in cancer development was believed to result from its ability to prevent cell death (thereby increasing cell number). We previously showed that bcl-2 deregulates G <sub>1</sub> /S checkpoint in MCF10A cells, which involves induction of cyclin D <sub>1</sub> associated kinase activity. Increasing evidence suggests that overexpression of cyclin D <sub>1</sub> contributes to the oncogenic transformation of cells <i>in vitro</i> and <i>in vivo</i> . In the present study, we examined the effects of bcl-2 overexpression on cyclin D <sub>1</sub> expression in human breast epithelial cells. Bcl-2 overexpression in MCF10A cells significantly induced expression of cyclin D <sub>1</sub> during the cell cycle as determined by an immunoblot analysis. We also show that bcl-2 induction of cyclin D <sub>1</sub> expression occurs at the transcriptional level as determined by an assay of cyclin D <sub>1</sub> promoter activity. Cyclin D <sub>1</sub> reporter activity was induced in a bcl-2 dependent manner in human breast carcinoma cell lines, MCF7 and BT549 as well as in MCF10A. Our previous and present studies suggest that bcl-2 may serve as an oncogene in the development of human breast cancer, which involves induction of cyclin D <sub>1</sub> expression.				
14. SUBJECT TERMS Breast Cancer			15. NUMBER OF PAGES 10	
			16. PRICE CODE	
17. SECURITY CLASSIFICATION OF REPORT Unclassified	18. SECURITY CLASSIFICATION OF THIS PAGE Unclassified	19. SECURITY CLASSIFICATION OF ABSTRACT Unclassified	20. LIMITATION OF ABSTRACT Unlimited	

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## Introduction

The proto-oncogene bcl-2 is frequently overexpressed in many human tumors including invasive breast cancers (1, 2). *In vitro* studies clearly demonstrate that the bcl-2 gene product prevents apoptosis following a variety of stimuli including radiation, hyperthermia, growth factor withdrawal and chemotherapeutic drugs. In transgenic mice, overexpression of bcl-2 under the immunoglobulin promoter induces follicular hyperplasia, suggesting a role for bcl-2 as an oncogene. However, high levels of bcl-2 expression have shown a positive correlation in clinicopathological studies such as tumor grade, and better response to hormone treatment and chemotherapy (3). The long-term goal of our studies is to investigate the *in vivo* functions of bcl-2 in human breast cancer development. To this end, we have used MCF10A and MCF10AT models.

MCF10A cell line was established in our institute without viral or chemical intervention from mortal diploid human breast epithelial cells (4). This cell line has been utilized by many laboratories to study sequential development of differentiated or malignant states of breast epithelial cells.

Dr. Miller (consultant in this application) and his colleagues have developed an *in vivo* model system to study human breast cancer progression (5-7); Whereas MCF10A cells do not survive *in vivo* in immune deficient mice, c-Ha-ras oncogene transfected MCF10A cells (MCF10AneoT) form small nodules in Nude/Beige mice which persist for at least one year and sporadically progress to carcinomas. Cell lines have been established in cultures from lesions representing 4 successive transplant generations (designated MCF10ATG1, MCF10ATG2, MCF10ATG3 and MCF10ATG4). With each generation, cells progress to high risk lesions resembling human proliferative breast disease. Thus, the MCF10AT model provides a setting in which the steps in the conversion of the breast ductal epithelial cell to malignant disease can be studied.

## Body of Report

During the 1997-1998 funding period, we have continued to investigate potential oncogenic activities of bcl-2 in breast cancer development.

## Methods

### Immunoblot analysis

MCF10A and bcl-2 overexpressing MCF10A clones (MCF10A bcl-2-6, MCF10A bcl-2-8, MCF10A bcl-2-30 and MCF10A bcl-2-40) were cultured as previously described (8). Cells were growth-arrested by culturing in serum-free medium for 48 hours at confluence. Cells were then treated with regular MCF10A medium containing 5% horse serum and growth factors to induce the cell cycle. At various times between 0-24 hr after serum-stimulation, whole-cell extracts were prepared using SDS lysate buffer. Protein concentrations were measured using bicinchoninic acid protein assay reagents (Pierce, Rockford, IL). Cell lysates (20 µg/lane) were denatured, subjected to SDS-PAGE analysis, and then electrophoretically transferred to a nitrocellulose membrane. Membranes were incubated with anti-cyclin D<sub>1</sub> (Ab2, Oncogene Research, MA), anti-bcl-2 or β-actin antibodies. Proteins were visualized using HRP-conjugated goat anti-mouse IgG (1:3000 dilution) and chemiluminescence reagent (Dupont, Boston MA 02118). The membranes were exposed to X-ray film from 1 to 15 minutes.

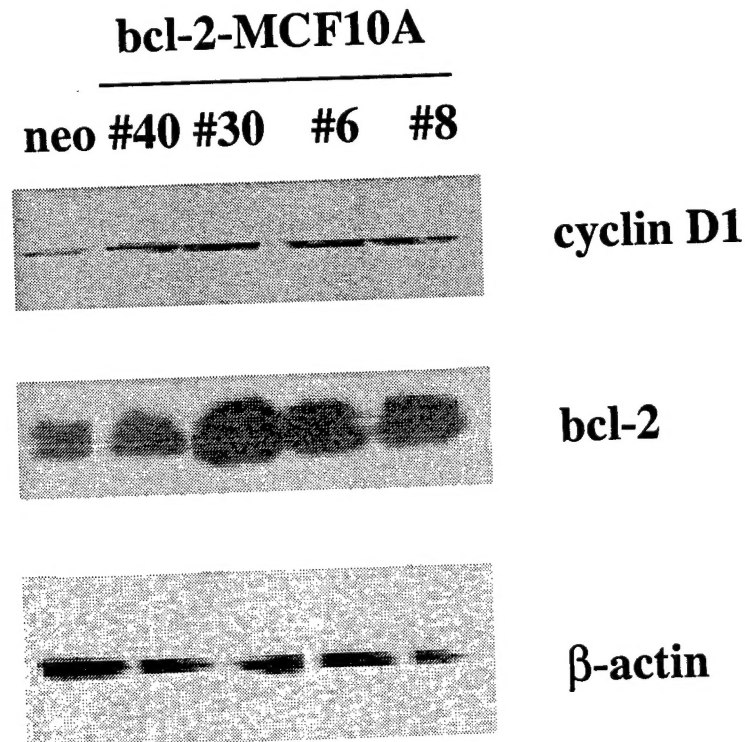
### Cyclin D<sub>1</sub> promoter activity assay

MCF10A, MCF7 and BT549 cells were transfected with -964 CD1 promoter, PCH110

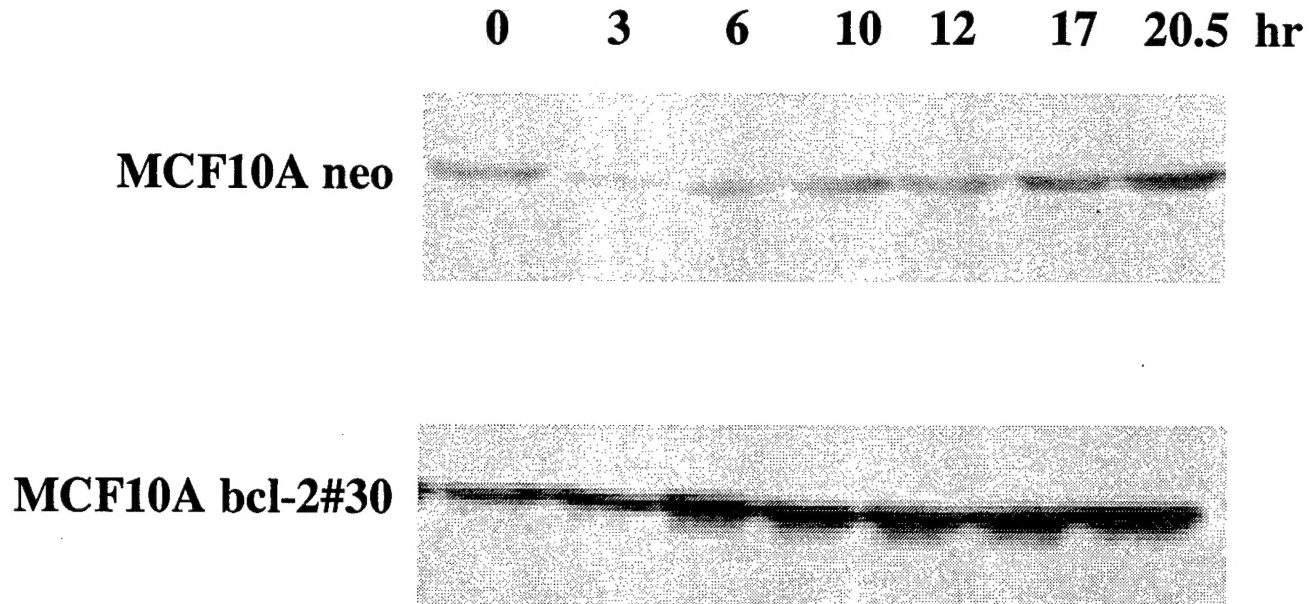
(MDVlacZ), and bcl-2 expression plasmid using FuGENE6 transfection reagent (Boehringer Mannheim). Fifty percent confluent cells in a 60-mm dish were transfected in 3 ml culture medium into which 500  $\mu$ l FuGENE6 reaction mixture was added. The FuGENE mixture was prepared as follows; 2  $\mu$ g of -964CD1 plasmid, 0.2  $\mu$ g of PCH110 and increasing amounts of bcl-2 expression plasmid were mixed in 250  $\mu$ l of serum free medium. The DNA solution was mixed with 250  $\mu$ l of serum free-medium which contains 50  $\mu$ l FuGENE6 reagent. The FuGENE6 mixture was incubated at room temperature for 30 minutes before transfection. Cells were harvested after 48 hours of transfection with 1X reporter lysis buffer. After mixing 20  $\mu$ l of cell lysate and 100  $\mu$ l luciferase substrate, the luciferase activity was measured using a luminometer. The  $\beta$ -galactosidase activity was measured by a chemiluminescent reporter assay (Tropix) to normalize transfection efficiencies. Protein concentration of the cell lysate was measured using the BCA reagent (Pierce).

## Results

We previously showed that bcl-2 deregulates G<sub>1</sub>/S check point through modulation of cyclin D<sub>1</sub>-associated kinase activity. In the present study, we examined whether bcl-2 regulates cyclin D<sub>1</sub> expression. When the levels of cyclin D<sub>1</sub> expression were examined in growing cells of MCF10A and bcl-2 overexpressing MCF10A cells, it appeared that bcl-2 induced expression of cyclin D<sub>1</sub> (Fig 1). To exclude the possibility that the differences in the levels of cyclin D<sub>1</sub> reflect the differences in the cell cycle distribution between the control and bcl-2 overexpressing cells, we examined the effects of bcl-2 overexpression on cyclin D<sub>1</sub> expression during the cell cycle. The control and bcl-2 overexpressing MCF10A cells were synchronized at G<sub>0</sub> by culturing in serum-free medium for 48 hours and the cell cycle was induced by culturing them in complete MCF10A culture medium. At various times between 0-24 hr after cell cycle induction, whole-cell extracts were prepared and the levels of cyclin D<sub>1</sub> were determined by immunoblot analysis. As shown in Fig. 2, the levels of cyclin D<sub>1</sub> expression were significantly higher throughout the cell cycle in bcl-2 overexpressing cells than in the control cells.



**Fig. 1. Bcl-2 induces cyclin D<sub>1</sub> expression in MCF10A cells.** The levels of cyclin D<sub>1</sub> (top panel) and bcl-2 (middle panel) proteins were determined by immunoblot analyses. Protein samples were prepared from neo-resistant vector transfected MCF10A (lane 1) and bcl-2 overexpressing MCF10A clones (lanes 2-5). To confirm the amount and quality of proteins loaded in each lane, the identical blot was probed with anti- $\beta$ -actin antibody (bottom panel).

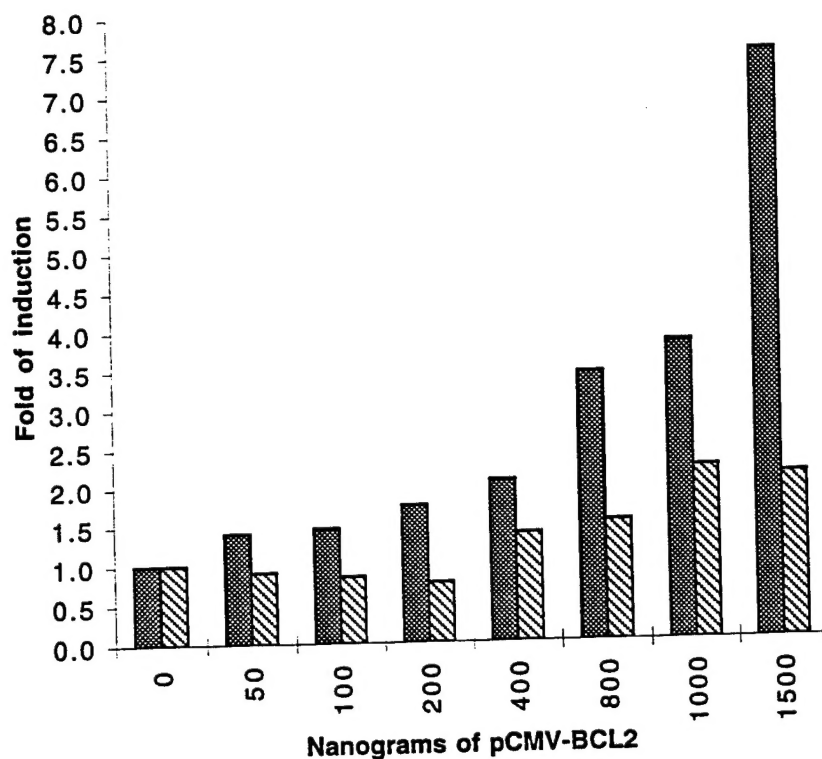


**Fig. 2. Bcl-2 induces cyclin D<sub>1</sub> expression during the cell cycle in MCF10A cells.** Confluent control (top panel) and bcl-2 overexpressing MCF10A clone 30 (bottom panel) were cultured in serum free DMEM/F12 medium for 48 hours. At various time points after serum treatment, cells were lysed using SDS sample buffer. Levels of cyclin D<sub>1</sub> protein were determined by immunoblot analyses.

We then examined whether bcl-2 induces transcription of cyclin D<sub>1</sub> in MCF10A cells. The effects of bcl-2 on cyclin D<sub>1</sub> promoter activity were determined by cotransfection experiments using -964CD1LUC plasmid (containing the human cyclin D<sub>1</sub> promoter linked to the luciferase reporter provided by Dr. Pestell, *see ref. 12*) and bcl-2 expression vector. As shown in Fig 3A, cyclin D<sub>1</sub> reporter activity was induced in a bcl-2 dependent manner. It was examined whether bcl-2 induction of cyclin D<sub>1</sub> transcription occurs in other human breast epithelial cells. To this end, we used human breast carcinoma cell lines, MCF7 and BT549. As shown in Fig 3A and B, bcl-2 effectively induced promoter activities of cyclin in these cell lines. These results show that bcl-2 induces expression of cyclin D<sub>1</sub> at the transcriptional level in human breast epithelial cells.

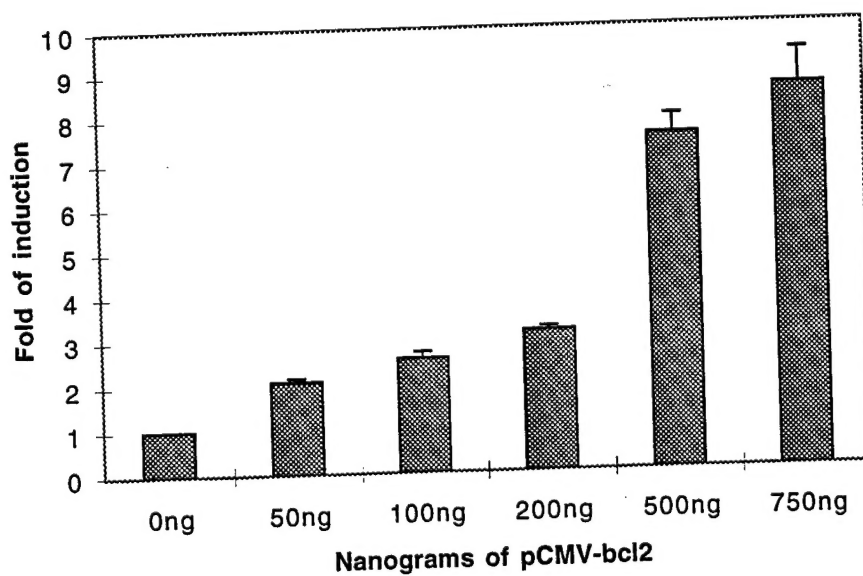
A.

### BCL-2 Induction of Cyclin-D1 Transcription



B.

### Bcl2 Induction of CyclinD1 Transcription





**Fig. 3. Bcl-2 induces cyclin D<sub>1</sub> promoter activity in human breast epithelial cells.**

A. -964CD1LUC reporter was transfected with increasing amounts of bcl-2 expression vector into MCF10A (hatched bar in panel A), BT549 (dotted bar in panel A) and MCF 7 (panel B). All experiments were performed in triplicates and the folds of induction represent the luciferase activity/LacZ activity/mg protein.

We are currently investigating the regions of the cyclin D<sub>1</sub> promoter responsible for transcriptional activation by bcl-2 overexpression using control and bcl-2 overexpressing cells. Our preliminary study showed that mutation of the AP1 binding site in the cyclin D<sub>1</sub> promoter does not have any effect on bcl-2 activation of cyclin D<sub>1</sub> promoter activity. However, deletion of the SP-1 binding site in -163 significantly reduced bcl-2 activation of cyclin D<sub>1</sub> promoter activity (data not shown). Further studies are in demand to determine the mechanisms by which bcl-2 activates cyclin D<sub>1</sub> promoter activity.

### Conclusion

Increasing evidence suggests that overexpression of cyclin D<sub>1</sub> contributes to the oncogenic transformation of cells *in vitro* and *in vivo* (9-11). Involvement of bcl-2 in cancer development was believed to result from its ability to prevent cell death (thereby increasing cell number). However, our previous and present studies suggest that bcl-2 may serve as an oncogene in the development of human breast cancer, which involves induction of cyclin D<sub>1</sub> expression. We now wish to determine the *in vivo* oncogenic role of bcl-2 using bcl-2 overexpressing MCF10ATG3B (we previously showed that bcl-2 overexpression in MCF10ATG3B induces a transformed phenotype as determined by a soft agar assay). Since the Career Development Award does not support any research expense, it is difficult to perform the proposed study which includes animal experiments.

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